## Genomewide Association Study of Severe Covid-19 with

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Citation Report

#	Article	IF	CITATIONS
1	Association of <scp>ABO</scp> blood group and secretor phenotype with severe <scp>COVID</scp> â€19. Transfusion, 2020, 60, 3067-3070.	0.8	32
2	Correlation of the two most frequent HLA haplotypes in the Italian population to the differential regional incidence of Covid-19. Journal of Translational Medicine, 2020, 18, 352.	1.8	86
3	Expression of ACE2, the SARS-CoV-2 Receptor, in Lung Tissue of Patients With Type 2 Diabetes. Diabetes, 2020, 69, 2691-2699.	0.3	55
4	Risk Variant for Severe COVIDâ€19 Inherited from Neanderthals. American Journal of Medical Genetics, Part A, 2020, 182, 2203-2204.	0.7	3
5	Genetics of COVID-19. Jornal De Pediatria, 2021, 97, 378-386.	0.9	17
6	Genetic Risk of Severe Covid-19. New England Journal of Medicine, 2020, 383, 1590-1591.	13.9	22
7	COVID-19 in Children: A Review and Parallels to Other Hyperinflammatory Syndromes. Frontiers in Pediatrics, 2020, 8, 593455.	0.9	16
8	On the genetics and immunopathogenesis of COVID-19. Clinical Immunology, 2020, 220, 108591.	1.4	32
9	Chemokine receptor gene polymorphisms and COVID-19: Could knowledge gained from HIV/AIDS be important?. Infection, Genetics and Evolution, 2020, 85, 104512.	1.0	16
10	<scp>SARS oV</scp> â€2 infection in <scp>India</scp> bucks the trend: Trained innate immunity?. American Journal of Human Biology, 2021, 33, e23504.	0.8	16
11	The influence of ABO blood groups on COVID-19 susceptibility and severity: A molecular hypothesis based on carbohydrate-carbohydrate interactions. Medical Hypotheses, 2020, 144, 110155.	0.8	42
12	IFN-γ is an independent risk factor associated with mortality in patients with moderate and severe COVID-19 infection. Virus Research, 2020, 289, 198171.	1.1	134
13	Genetic Determinants of Antibody-Mediated Immune Responses to Infectious Diseases Agents: A Genome-Wide and HLA Association Study. Open Forum Infectious Diseases, 2020, 7, ofaa450.	0.4	12
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16	Understand variability of COVID-19 through population and tissue variations in expression of SARS-CoV-2 host genes. Informatics in Medicine Unlocked, 2020, 21, 100443.	1.9	24
17	Transcriptional and proteomic insights into the host response in fatal COVID-19 cases. Proceedings of the United States of America, 2020, 117, 28336-28343.	3.3	149
18	The antibody response to the glycan αâ€Gal correlates with COVIDâ€19 disease symptoms. Journal of Medical Virology, 2021, 93, 2065-2075.	2.5	25

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19	Possible role of HLA class-I genotype in SARS-CoV-2 infection and progression: A pilot study in a cohort of Covid-19 Spanish patients. Clinical Immunology, 2020, 219, 108572.	1.4	76
20	Evaluation of a genetic risk score for severity of COVID-19 using human chromosomal-scale length variation. Human Genomics, 2020, 14, 36.	1.4	8
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24	The association of ABO blood group with indices of disease severity and multiorgan dysfunction in COVID-19. Blood Advances, 2020, 4, 4981-4989.	2.5	128
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56	Associations between blood type and COVID-19 infection, intubation, and death. Nature Communications, 2020, 11, 5761.	5.8	309
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	<ul> <li>Hematology, 2020, 100, 3081-3082.</li> <li>Genetic architecture of host proteins involved in SARS-CoV-2 infection. Nature Communications, 2020, 11, 6397.</li> <li>No impact of cancer and plague-relevant <i>FPR1</i> polymorphisms on COVID-19. Oncolmmunology,</li> </ul>	5.8	71
122	<ul> <li>Hematology, 2020, 100, 3081-3082.</li> <li>Genetic architecture of host proteins involved in SARS-CoV-2 infection. Nature Communications, 2020, 11, 6397.</li> <li>No impact of cancer and plague-relevant <i>FPR1</i> polymorphisms on COVID-19. Oncolmmunology, 2020, 9, 1857112.</li> <li>Temporal and spatial heterogeneity of host response to SARS-CoV-2 pulmonary infection. Nature</li> </ul>	5.8 2.1	71 4
122 123	Hematology, 2020, 100, 3081-3082.         Genetic architecture of host proteins involved in SARS-CoV-2 infection. Nature Communications, 2020, 11, 6397.         No impact of cancer and plague-relevant <i>FPR1</i> polymorphisms on COVID-19. Oncolmmunology, 2020, 9, 1857112.         Temporal and spatial heterogeneity of host response to SARS-CoV-2 pulmonary infection. Nature Communications, 2020, 11, 6319.         Comparison of clinical, para-clinical and laboratory findings in survived and deceased patients with COVID-19: diagnostic role of inflammatory indications in determining the severity of illness. BMC	5.8 2.1 5.8	71 4 203
122 123 124	Hematology, 2020, 100, 3081-3082.         Genetic architecture of host proteins involved in SARS-CoV-2 infection. Nature Communications, 2020, 11, 6397.         No impact of cancer and plague-relevant <i>FPR1</i> polymorphisms on COVID-19. Oncolmmunology, 2020, 9, 1857112.         Temporal and spatial heterogeneity of host response to SARS-CoV-2 pulmonary infection. Nature Communications, 2020, 11, 6319.         Comparison of clinical, para-clinical and laboratory findings in survived and deceased patients with COVID-19: diagnostic role of inflammatory indications in determining the severity of illness. BMC Infectious Diseases, 2020, 20, 869.         Prognostic Genetic Markers for Thrombosis in COVID-19 Patients: A Focused Analysis on D-Dimer,	5.8 2.1 5.8 1.3	71 4 203 36

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